AMENDMENTS TO THE CLAIMS

This listing of claims will replace all prior versions and listings of claims in the application.

Listing of Claims

- 1. (Currently amended) A composition comprising a protein in crystalline form wherein at least a portion of the protein has at least 90% identity with consists of SEQ. ID No. 1.3.
- 2-3. (Cancelled)
- 4. (Currently amended) A composition according to claim 1 wherein the protein crystal diffracts X-rays for a determination of structure coordinates to a resolution greater than having a value that is less than or equal to 3.0 Angstroms.
- 5. (Original) A composition according to claim 1 wherein the protein crystal has a crystal lattice in a P4₁22 space group.
- 6. (Original) A composition according to claim 1 wherein the protein crystal has a crystal lattice having unit cell dimensions, +/- 5%, of a=b= 85.159Å and c=152.18Å.
- 7-8. (Cancelled)
- 9. (Currently amended) A method for forming a crystal of a protein comprising:

forming a crystallization volume comprising: a precipitant solution and a protein wherein at least a portion of the protein has at least 90% identity with that consists of SEQ. ID No.-1_3; and

storing the crystallization volume under conditions suitable for crystal-formation of the <u>a</u> protein <u>crystal</u>.

10-11. (Cancelled)

- 12. (Currently amended) A method according to claim 9 wherein the <u>a</u> protein crystal <u>is</u> formed that diffracts X-rays for a determination of structure coordinates to a resolution greater than having a value that is less than or equal to 3.0 Angstroms.
- 13. (Currently amended) A method according to claim 9 wherein the <u>a</u> protein crystal <u>is</u> formed that has a crystal lattice in a P4₁22 space group.
- 14. (Currently amended) A method according to claim 9 wherein the <u>a</u> protein crystal <u>is</u> formed that has a crystal lattice having unit cell dimensions, +/- 5%, of a=b= 85.159Å and c=152.18Å.
- 15. (Currently amended) A method according to claim 9, wherein a protein crystal is formed, the method further comprising diffracting the protein crystal to produce a diffraction pattern and solving the structure of the protein from the diffraction pattern.
- 16-17. (Cancelled)
- 18. (Currently amended) A composition comprising an isolated a protein consisting of SEQ. ID No. 3.
- 19-26. (Cancelled)
- 27. (New) The method according to claim 15 further comprising: performing rational drug design using the solved structure; and identifying an entity that associates with the protein.
- 28. (New) The method according to claim 27 further comprising selecting one or more entities based on the rational drug design and contacting the selected entities with the protein.
- 29. (New) The method according to claim 28 further comprising measuring an activity of the protein when contacted with the one or more entities.